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Abstract

We investigated the effect of -174 G/C single-nucleotide polymorphism in the promoter region of the *IL6* gene on plasma IL-6 levels and muscle strength, and the relationship between IL-6 levels and muscle strength in elderly women. The sample consisted of 199 elderly residents (73.0 ± 7.8 years old) from rest homes and the community in Belo Horizonte, MG, Brazil. -174 G/C polymorphism was determined by direct sequencing of the product by PCR, and plasma IL-6 concentrations were measured by ELISA. Muscle strength in the knee joint was evaluated using a Biodex System 3 Pro[®] isokinetic dynamometer. ANCOVA was used to determine the effect of polymorphism on IL-6 levels and muscle strength, and the Pearson correlation coefficient to assess the relationship between IL-6 levels and muscle strength. -174 G/C polymorphism was associated with the plasma IL-6 levels of elderly women ($P < 0.01$) since homozygotes for the G allele showed high IL-6 levels (GG 3.85 pg/mL, GC + CC 2.13 pg/mL). There was no association of polymorphism on muscle strength ($P > 0.05$). No association was found between IL-6 levels and knee extensor muscle ($r = 0.087$, $P = 0.306$) or flexor ($r = -0.011$, $P = 0.894$) strength. An interaction between -174 G/C polymorphism and housing conditions of the sample of elderly women was identified, with the effect of genotype on IL-6 levels being higher in the institutionalized elderly. These results support the evidence that -174 G/C polymorphism of the *IL6* gene associates with individual variability of plasma IL-6 levels in elderly women.

Key words: Interleukin-6; Elderly; Muscular strength; -174 G/C polymorphism

Introduction

The aging process is associated with subliminal chronic inflammatory activity, with the occurrence of an idiopathic elevation of serum levels of inflammatory mediators such as interleukin-1 (IL-1), tumor necrosis factor- α (TNF- α), interleukin-6 (IL-6), and acute phase proteins such as C-reactive protein (CRP), among other mediators (1,2). This subliminal state of inflammation has been linked to the pathological process of various age-related diseases and to the increased disability and mortality of the elderly (1-4).

IL-6 is a multifunctional pleiotropic cytokine with recognized pro- and anti-inflammatory action. It plays an important

role in the homeostasis of the immune and neuroendocrine systems, as well as in the balance of proinflammatory/anti-inflammatory pathways and in response to stress (1,2,5). There is evidence that the concentration of IL-6 increases in the supernatant of peripheral blood mononuclear cell cultures and in the plasma of elderly individuals (1,6,7). Increased plasma IL-6 levels are associated with the presence of cardiovascular and rheumatic diseases, sarcopenia, functional decline, and mortality in the elderly (1,7,8).

The expression of IL-6 is related to its allelic variant (9). It is known that the *IL6* gene has about 50 single-nucleotide

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polymorphisms (SNP) in its promoter region, like the variants -597 G/A, -572 G/C, -373 A/T, and -174 G/C, among others (10-12). Several studies have questioned the functional role of the -174 G/C SNP in the production of IL-6 both *in vivo* (13-15) and *in vitro* (10-12). The change of guanine bases to cytosine (G → C) at position -174 seems to affect the transcription of the *IL6* gene and therefore the plasma levels of this cytokine in young (10,12), elderly (12,13,15) and centenarian individuals (13,16). However, the results of studies regarding the activity and effects of polymorphisms on plasma levels of these mediators and susceptibility to different diseases are contradictory.

Some evidence suggests that genetic factors may influence the physical functioning of elderly people since the hereditary factor for muscle strength may vary from 36 to 65% (17-19). However, there is little information about specific genes that may contribute to the physical disability related to aging. In this respect, the genes involved in regulating the inflammatory process, which increases with age, may contribute to individual variation in plasma cytokine levels and muscle strength in the elderly. However, despite evidence pointing at the action of IL-6 as one of the factors related to the reduction of muscle mass and muscle strength (sarcopenia) (1,20-22) and the importance of genetic variation in the regulation of this cytokine, studies investigating the impact of the -174 G/C polymorphism of the *IL6* gene on the variables of muscle performance are scarce.

The objective of the present study was to investigate the effect of -174 G/C SNP in the promoter region of the *IL6* gene on plasma IL-6 levels and muscle strength in elderly women. The association between plasma IL-6 levels and muscle strength was also analyzed.

Material and Methods

This was a cross-sectional observational study with a convenience sample consisting of elderly female residents of homes for the aged and in the community, in the city of Belo Horizonte. The study was approved by the Ethics Committee of Universidade Federal de Minas Gerais (ETIC 322/07) and all volunteers gave written informed consent to participate, according to the principles of the Helsinki Declaration.

Sample

The sample was a convenience selection, whereby the participants were recruited by consecutive recruiting visits to homes for the aged, health centers and senior citizen social groups and by announcements in local homes for the aged and in the community. Exclusion criteria were: elderly people with acute phase inflammatory disease, using drugs that act on the immune system, presenting cognitive impairment detectable by the Mini-Mental State Examination (23), amputation or lower limb fracture in the last 6 months, and the presence of neurological sequelae.

For sampling in the community, 455 potential volunteers were initially contacted, and 153 elderly subjects were selected on the basis of the inclusion and exclusion criteria. Of these, 8 dropped out: 4 because of health problems (cataract surgery, erysipelas, deep vein thrombosis, and cardiac abnormalities) and 4 because of transportation difficulties to reach the site of assessment. The total number of elderly subjects was 145, who were from the community. Due to the high prevalence of cognitive disorders, sickness in the acute phase, use of medications acting on the immune system and refusal of some homes for the aged to participate, the sample from homes for the aged consisted of only 57 elderly women. Therefore, with the elderly from the community and the homes for the aged, a total of 202 participants were selected for the study.

To characterize the sample, the sociodemographic data (age, marital status, education, and personal income) and information on the clinical conditions of the elderly (comorbidities, number and type of medications, body mass index (BMI), presence of pain, walking capacity, and occurrence of falls in the previous year) were obtained using a structured questionnaire and a clinical assessment, carried out by trained personnel.

Muscle strength

Muscle strength was assessed using a Biodex® System 3 Pro isokinetic dynamometer (Biodex Medical Systems Inc., USA). The muscle groups assessed were the knee joint extensors and flexors of the dominant limb. Work was standardized according to body mass, selected as a parameter of muscle performance, and was obtained by means of concentric contractions at an angular velocity of 60°/s, with five repetitions. All procedures were performed according to the assessment protocol suggested by the manufacturer, such as positioning of the volunteer, calibration, correction for gravity, familiarization, and strong verbal encouragement (24). Considering the difficulties in selecting and transporting the institutionalized elderly to the place of assessment, muscle strength was measured only in the elderly from the community.

IL-6 levels

Five milliliters of blood was collected from the ulna vein of the participants into Vacutainers with citrate by a qualified healthcare professional. The blood was then centrifuged and the plasma was removed in a sterile environment and stored in Eppendorf tubes in a freezer at -70°C for 2 years.

Plasma IL-6 concentrations were measured by ELISA using high sensitivity kits from Quantikine® HS, R&D Systems, USA, according to manufacturer recommendations. Measurements were made in duplicate and results are reported as means ± SD. The intra-assay coefficient of variation was 6.9-7.4%, the interassay coefficient of variation was 9.6-6.5%, and sensitivity was 0.016-0.110 pg/mL (mean: 0.039 pg/mL). Data are reported as picograms per

mL (pg/mL). Absorbance was measured using a single microplate reader at 490 nm with length correction at 650 nm (Microplate Reader Asys, Austria).

Identification of genotype

Total DNA was isolated from peripheral plasma samples using the QIAamp DNA Mini-Kit (Qiagen, USA) according to manufacturer recommendations. The -174 G/C polymorphism in the promoter region of the *IL6* gene (rs1800795) was determined by direct sequencing of the product of a polymerase chain reaction (PCR). A 628-bp region of the *IL6* gene containing the polymorphic point was amplified using a specific pair of primers: 5'-GAACACAGAAGAAGCTCAGATGACTGG-3' (sense) and 5'-AGGAGTTCATAGCTGGGCTCCTGGAG-3' (antisense). The reactions were performed in tubes containing 100 ng DNA, 10 mM Tris-HCl, pH 9.2, 25 mM KCl, 1.5 mM MgCl₂, 0.2 mM dNTP, 20 pmol of each primer, 0.5 µg purified albumin (purified chicken albumin) and one unit of *Taq* polymerase DNA (Phoneutria, Brazil) in a final volume of 50 µL. After 1 min of initial heating at 80°C and an initial denaturation for 2 min at 94°C, amplification was performed for 36 cycles of 40 s at 94°C, 45 s at 64°C, and 50 s at 72°C, followed by 5 min at 72°C. A sample (100 ng) of each PCR product was sequenced directly with an ABI PRISM 3700 DNA analyzer (Applied Biosystems, USA) using the DYEnamic ET Terminator Cycle Sequencing Kit (GE Healthcare, UK) along with 2 µM of the 5'-GCCTCAGAGACATCTCCAGTCC-3' primer in the reaction tube. Sequencing conditions using a PE 9700 thermal cycler (Perkin Elmer, USA) were as follows: 30 cycles of 20 s at 95°C, 15 s at 50°C, and 1 min at 60°C. Each sequence obtained was examined using the Staden software package (MRC, UK) and confirmed by visual inspection.

Statistical analysis

The description of the subject sample was done by measuring central tendency (means ± SD) and frequency of the socio-economic and clinical variables. Violation of Hardy-Weinberg equilibrium was tested by the chi-square test, taking $P < 0.05$.

The effect of polymorphism on IL-6 levels and muscle strength in the elderly women was investigated by analysis of covariance (ANCOVA). Considering the heterogeneity of the sample and the known association of some variables with plasma IL-6 indexes and muscle strength, a covariance test was performed in which IL-6 levels were considered as dependent variable regarding genotype, housing conditions (institutionalized or living in the

community) and the presence of osteoarthritis constituted the fixed factors (independent). Age and BMI were used as covariants.

The Pearson correlation coefficient (r) was used to assess the correlation between plasma IL-6 levels and muscle strength. Data normality was checked by the Kolmogorov-Smirnov test and homogeneity of variances by the Levene test. The level of significance was set at $\alpha = 0.05$ in all analyses. Analyses were performed using the Statistical Package for Social Sciences (SPSS) for Windows (version 15.0).

Results

Of the 202 elderly subjects from the community and homes for the aged that met the inclusion criteria, 199 participated in the study, 55 of them residents of homes for the aged and 144 residents of the community. Three elderly women (2 institutionalized and 1 from the community) were excluded because it was not possible to obtain their genotype for the -174 G/C polymorphism of the *IL6* gene. The sociodemographic and clinical characteristics of the total sample are presented in Table 1.

The genotypes were consistent with Hardy-Weinberg equilibrium proportions (58% GG, 39% GC, and 3% CC; $P > 0.05$). The frequency of genotypes for the -174 G/C polymorphism and the plasma levels of IL-6 are presented in Table 2. For analysis, the genotypes were

Table 1. Sociodemographic and clinical characteristics of the elderly women assessed.

Variable	Home for the aged N = 55 (28%)	Community N = 144 (72%)	Total sample N = 199
Age (means ± SD)	77.4 ± 9.5	71.3 ± 6.2	73.0 ± 7.8
Educational level			
No schooling	13 (24%)	13 (9%)	26 (13%)
1 to 7 years of schooling	31 (56%)	63 (11%)	94 (47%)
8 or more years of schooling	11 (20%)	68 (47%)	79 (40%)
Marital status			
Single	35 (64%)	6 (4%)	41 (21%)
Married	0	59 (41%)	59 (30%)
Widow	17 (31%)	57 (40%)	74 (37%)
Divorced	3 (5%)	22 (15%)	25 (13%)
Income			
<2 minimum wages*	51 (93%)	46 (32%)	96 (49%)
2-3.99 minimum wages	4 (7%)	51 (35%)	55 (28%)
≥4 minimum wages	0	46 (32%)	46 (23%)
BMI (kg/m ²)	25.3 ± 5.5	28.9 ± 4.7	28.0 ± 5.2
Osteoarthritis	26 (47%)	113 (79%)	139 (70%)
HBP	27 (49%)	91 (63%)	118 (59%)

*Minimum wage for 2010 = R\$510.00 (US\$238.31 to US\$305.38). The categorical variables (educational level, marital status and income) are reported as frequency and percentage. BMI = body mass index; HBP = high blood pressure.

grouped as GG vs GC + CC.

The difference in mean plasma IL-6 levels was significant between the groups of genotype GG vs GC + CC, with the elderly women with the GG genotype having higher levels of this cytokine when compared with carriers of the C allele (ANCOVA). The effect of polymorphism on IL-6 levels was significant even under the influence of, and interaction between, the housing condition variables (homes for the aged or community), and the presence of osteoarthritis and BMI were controlled ($F = 10.99$; $P < 0.01$).

In the analysis of variance of IL-6 levels attributed to genotype and housing condition factors, the polymorphism showed a higher relative effect of IL-6 levels (partial $\eta^2 = 0.058$), which, however, was lower than the effect of the housing conditions on this mediator (partial $\eta^2 = 0.189$). Moreover, an interaction between residing status and -174 G/C polymorphism ($F = 5.84$; $P = 0.02$) was observed, with the effect of genotype on IL-6 levels being higher in the institutionalized elderly compared to the elderly women from the community. Comparison of IL-6 levels between groups of genotypes (GG vs GC + CC) revealed more marked levels in the institutionalized elderly women compared to those living in the community ($P < 0.05$; Table 3).

To analyze the effect of polymorphism on muscle strength, 139 of 144 elderly women from the community were assessed using an isokinetic dynamometer, and 5 dropped out during this phase of the study. No difference was observed in the muscle strength of knee joint extensors and flexors, assessed according to work/body mass in the elderly from the community, between groups of genotypes (GG vs GC + CC) ($P > 0.05$). Similarly, no correlation was found between plasma IL-6 levels and knee extensor muscle.

Discussion

Results obtained for the sociodemographic profile and the clinical population studied were consistent with literature report regarding low education and low income. The sample presented a tendency toward excess weight according to Lipschitz (25) cutoff points for the elderly, and the prevalence of osteoarthritis was high.

Analysis of covariance showed that the -174 G/C polymorphism affected the plasma IL-6 levels of elderly women regardless of whether they resided in the community or were institutionalized ($P < 0.01$). Higher IL-6 levels were observed in the group of elderly women with the GG genotype, as also observed in another recent Brazilian study (26). This effect was significant even when factors related to IL-6 levels were controlled, such as the following variables: housing conditions (homes for the aged or community), presence of osteoarthritis, age and BMI. However, the housing condition variable for the elderly women strongly affected the plasma IL-6 levels, with institutionalized women presenting comparatively higher IL-6 levels than women

Table 2. Distribution of genotypes and plasma IL-6 levels in elderly women living in homes for the aged or in the community.

Variable	Home for the aged	Community	Total sample
GG	34 (62%)	82 (57%)	116 (58%)
GC	20 (36%)	58 (40%)	78 (39%)
CC	1 (2%)	4 (3%)	5 (3%)
Plasma IL-6 (pg/mL)	6.37 ± 6.31	1.87 ± 2.49	3.13 ± 4.40

Data are reported as means ± SD. IL-6 = interleukin-6.

Table 3. Comparison of IL-6 levels between groups of genotypes GG vs GC + CC for IL6 gene -174 G/C polymorphism.

Variable	Plasma IL-6 (pg/mL)		ANCOVA	
	Home for the aged	Community	F	P
GG	7.74 ± 6.5	2.24 ± 2.3	5.84	0.02*
GC + CC	4.17 ± 3.4	1.36 ± 1.3		

Data are reported as means ± SD. Comparison of IL-6 levels between groups of genotypes (GG vs GC + CC) revealed more marked levels in elderly women living in homes for the aged compared to those living in the community ($F = 5.84$; $P = 0.02$). IL-6 = interleukin-6.

from the community.

The institutionalized elderly women usually represent a distinct population: they have more associated diseases, comorbidities, a higher prevalence of depression and difficulty to perform daily tasks. Moreover, they live without a multigenerational family and social contact and are submitted to a routine and to interventions that are peculiar to the institution where they live. Research has documented the impact of the social support network on the dysregulation of physiological parameters, where there is a lack of this association with morbidity and mortality (27,28). Growing evidence points to a relationship between social conditions and adverse environmental conditions and high levels of inflammatory mediators (27,28). Due to this, the living conditions of institutionalized elderly women can contribute to a chronic condition of biopsychosocial stress, which in turn can influence the plasma levels of proinflammatory mediators. These factors could explain the impact of institutionalization on the increased plasma IL-6 levels in the sample studied.

In the present study, a significant interaction effect between genotype and housing conditions of the elderly was also observed. The institutionalized elderly showed higher levels of plasma IL-6 compared to those living in the community, indicating that the effect of genotype on

IL-6 levels appears to be greater in the institutionalized elderly. Recent studies have shown that the control of the expression of IL-6 is extremely complex, being regulated by different mechanisms and factors (9,11,29). The promoter region of *IL6* genes behaves as a sophisticated biosensor regarding the homeostasis of the immune system, hormonal changes and different environmental factors such as diet and stress conditions as determined by lifestyle, social and psycho-emotional issues (30). In this context, the condition of institutionalization of elderly women could lead to imbalances of a nutritional and psychosomatic nature with inflammatory repercussions, causing the effect of a genotype prone to a higher production of cytokines.

However, even being influenced by environmental factors, the effect of -174 G/C polymorphism on IL-6 levels was significant for both the institutionalized and community-living elderly women. Indeed GG homozygous individuals showed higher levels of IL-6 even in distinct social and environmental conditions, providing evidence that this genotype interfered significantly with plasma IL-6 levels in the sample studied. These findings provide further support for literature data showing that the -174 GG genotype is associated with greater production of IL-6 both *in vitro* (10,12) and *in vivo* (12,13), but they differ from reports showing that this genetic influence is dependent on gender (12). Olivieri et al. (12) analyzed the effect of -174 G/C polymorphism on the production of IL-6 *in vitro* and *in vivo* in a sample of 62 individuals aged 29 to 93 years. A significant association between -174 G/C polymorphism and the production of IL-6 was observed both *in vitro* and *in vivo*, with GG individuals presenting higher IL-6 levels than CC + GC subjects, with a significant association only for males. These results agree with those reported by Bonafé et al. (13). However, information regarding gender specificities is limited, but there are indications of possible interactions between genes and sex hormones (31).

Other investigators, studying the association between -174 G/C polymorphism and specific diseases and longevity, have obtained contradictory results regarding genotypes (14,16,32,33). Rea et al. (34) evaluated changes in the frequency of GG homozygotes for polymorphism at position -174 G/C of the gene for IL-6 with aging and observed a 10% decline in the frequency of this genotype with advancing age, with homozygotes for the C allele presenting higher IL-6 levels. A study consistent with these results was conducted on individuals older than 80 years, with follow-up for a period of 6 years, where the C allele was associated with high levels of plasma IL-6 in octogenarians (14).

An important question should be considered regarding the conflicting results in the literature. The aging process and the health conditions related to age are complex phenotypes, influenced by the interaction between genes and different environmental factors. The different results observed among studies were possibly due to the interactions between

lifestyle and genetic factors, along with cultural and ethnic differences across different countries.

There is little research on the impact of the -174 G/C polymorphism of the *IL6* gene on muscle performance variables. In the present study, no effect of -174 G/C polymorphism of the *IL6* gene was observed on muscle strength of knee extensors and flexors, assessed by the work/body mass of elderly women in the community. These results support other results reported in the literature (35,36). Roth et al. (35) investigated the association between -174 G/C polymorphism of the *IL6* gene and muscle mass and strength in 242 individuals (110 men and 132 women) aged 21 to 92 years. These investigators found an association of this polymorphism with muscle mass, but not with muscle strength (35). Walston et al. (36) investigated the association of different alleles of the *IL6* gene with variables such as plasma IL-6 levels, muscle strength and fragility, in two cohorts of elderly subjects. No relationship was demonstrable between plasma IL-6 levels, muscular strength (handgrip and isometric strength of knee extensors and hip flexors) and weakness and the polymorphisms or the haplotypes investigated for the *IL6* gene, including the -174 G/C polymorphism.

However, these findings should be interpreted with caution since an approach focused on the polymorphism of a single inflammatory mediator is limited. Other mediators besides IL-6 play an important catabolic role in muscle and, likewise, their levels are influenced by functional polymorphisms in the promoter region of their genes, such as TNF- α . Another point to be considered is that possible interactions between genes of these cytokines may contribute to the variability in muscle performance observed with age, and mask the separate effects of a single polymorphism. This requires a more detailed investigation of the complex interactions between polymorphisms of different genes and their influence on specific phenotypes, such as muscle strength.

Laboratory and clinical studies have shown that high levels of IL-6 are associated with a reduction of muscle mass and strength in the elderly (8,37,38). The mechanism by which these cytokines contribute to muscle alteration is not known, but involves its catabolic action on muscle, with a reduction of myofibril protein being related to sarcopenia (8). However, in the present sample of community-living elderly women, no correlation was observed between muscle strength and plasma IL-6 levels. This result may be explained by the fact that the plasma levels found in the elderly assessed here (1.87 ± 2.49 pg/mL) did not reach the threshold necessary to influence muscle performance. In a cohort of 627 elderly women presenting moderate to severe disabilities, Ferrucci et al. (37) found that those with plasma IL-6 levels in the highest tertiles (IL-6 >3.1 pg/mL) showed a greater reduction of knee extensor muscle strength and also of the ability to perform daily activities.

Some limitations of the present study should be consid-

ered. The fact that it was a convenience sample does not permit to generalize the results, limiting its external validity. Another point to be considered is that the level of physical activity of the elderly women was not recorded. There is evidence that physical activity has an anti-inflammatory effect, with a reduction in plasma levels of inflammatory mediators, so that physically active individuals present lower levels of plasma IL-6. IL-6 is released during physical exercise, produced by the muscle (myocyte) and is accompanied by the synthesis of anti-inflammatory cytokines such as IL-1ra, a soluble TNF- α receptor (sTNFR) and IL-10. Moreover, physical activity has been shown to be an effective strategy for the prevention and treatment of sarcopenia (39,40). Thus, the practice of physical activity, if carried out by participants, may have influenced the results found regarding the effect of genotypes on muscle strength, and may have contributed to the lower levels of IL-6 found in the elderly women from the community.

The results of this study support the evidence that the

-174 G/C polymorphism of the *IL6* gene influences the levels of plasma IL-6 in elderly residents of homes for the aged and in the community. This effect persisted even when the influence and interactions between factors known to interfere with the production of IL-6 were controlled. Moreover, an interaction between -174 G/C polymorphism and the housing conditions of the women was identified, suggesting that environmental factors may modulate the effects of genotypes on the production of IL-6. No effect of polymorphism was observed on the muscle strength of the extensors and flexors of the knee joint and no association was found between plasma IL-6 levels and muscle strength in elderly women from the community.

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